AMENDMENT

In the claims:

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1. (currently amended) A method for creating automated inferences of physico-chemical

relationships, comprising:

(a) extracting a database record from a structured literature database;

(b) parsing the database record to extract one or more individual information fields,

wherein the one or more individual information fields include a set of chemical or biological

molecule names;

(c) filtering the extracted set of chemical or biological molecule names to create a

filtered set of chemical or biological molecules names;

(d) determining whether a chemical or biological molecule name from the filtered set

has been stored in an inference database,

and if not,

storing the chemical or biological name in the inference database,

and setting a co-occurrence count to a starting value for each pair of names including the

chemical or biological name and other names from the filtered set that the chemical or

biological name co-occurs with;

and if so,

incrementing co-occurrence counts for each pair of chemical or biological

names including the chemical or biological name;

(e) repeating steps (a)-(d) for unique database records in the structured literature

database;

McDonnell Boehnen Hulbert & Berghoff LLP 300 S. Wacker Drive Chicago, IL 60606 (f) optionally constructing a connection network using a plurality of database records

from the inference database including co-occurrence counts;

(fg) applying one or more analysis methods directly to database records in the

inference database or to the optional connection network to determine metrics

representative of possible inferences of physico-chemical relationships between chemical or

biological molecules; and

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(gh) generating automatically a plurality of inferences regarding physico-chemical

relationships between chemical or biological molecules using the metrics resulting results

from the one or more analysis methods.

2. (currently amended) The method of Claim 1 further comprising providing a

computer readable medium and storing having stored therein instructions for causing a

processor to execute the steps of the method.

3. (original) The method of Claim 1 wherein extracting step includes extracting a

plurality of database records with a pre-determined database record structure.

4. (currently amended) The method of Claim 3 wherein the extracting step includes

extracting a database record with a pre-determined structure from a database containing

indexed references. Medline, PubMed, Biological Abstracts or Science Citation Index

databases.

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5. (currently amended) The method of Claim 1 wherein the parsing step includes

parsing the database record to extract a record an information field indicating two or more

chemical or biological molecule names used in an experiment recorded in the database

record.

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6. (original) The method of Claim 1 wherein the filtering step includes filtering the

chemical or biological molecule names against a list of trivial chemical or biological molecule

names to be ignored.

7. (currently amended) The method of Claim 1 further comprising wherein the step of

optionally constructing a connection network includes constructing a connection network

including a plurality of nodes representing a plurality of chemical or biological molecules

names and a plurality of arcs connecting the plurality of nodes, wherein the plurality of arcs

represent co-occurrences between chemical or biological molecules.

8. (original) The method of Claim 1 wherein the applying step includes applying

statistical analysis methods to co-occurrence counts stored in the inference database.

9. (original) The method of Claim 1 wherein the generating step includes generating

automatically inferences for physico-chemical interactions between chemical or biological

molecules using the co-occurrence counts stored in the inference database.

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10. (currently amended) The method of Claim 9 wherein the physico-chemical

interactions between chemical or biological molecules include physico-chemical interactions

for chemical or biological molecules for biological cells.

11. (original) The method of the Claim 1 wherein the chemical or biological molecule

names include natural or synthetic chemical compound or chemical molecule names or

natural or synthetic biological molecule or biological compound names.

12. (original) The method of Claim 1 further comprising storing the plurality of

inferences in the inference database.

13. (original) The method of Claim 1 further comprising applying subsequent analysis

methods to the connection network to reject trivial inference associations.

14. (original) The method of Claim 13 wherein the subsequent analysis methods

include assigning derived numerical values to arcs in the connection network based on co-

occurrence counts, assigning derived numerical values to arcs in the connection network

based on analysis of a temporal pattern of an inference association's co-occurrence count as

a function of another variable, conducting a mutual information analysis, or conducting a

Citation analysis.

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15. (original) The method of Claim 1 wherein the step incrementing step includes incrementing a plurality of co-occurrence counts for pairs of chemical or biological molecule names in the filtered set.